

## SEARCH REQUEST FORM

Requestor's Name: \_\_\_\_\_ Serial Number: \_\_\_\_\_

Date: \_\_\_\_\_ Phone: \_\_\_\_\_ Art Unit: \_\_\_\_\_

### Search Topic:

Please write a detailed statement of search topic. Describe specifically as possible the subject matter to be searched. Define any terms that may have a special meaning. Give examples or relevant citations, authors, keywords, etc., if known. For sequences, please attach a copy of the sequence. You may include a copy of the broadest and/or most relevant claim(s).

### STAFF USE ONLY

Date completed: \_\_\_\_\_  
Searcher: Beverly C 4994  
Terminal time: \_\_\_\_\_  
Elapsed time: \_\_\_\_\_  
CPU time: \_\_\_\_\_  
Total time: \_\_\_\_\_  
Number of Searches: \_\_\_\_\_  
Number of Databases: 1

#### Search Site

\_\_\_\_\_ STIC  
\_\_\_\_\_ CM-1  
\_\_\_\_\_ Pre-S

#### Type of Search

\_\_\_\_\_ N.A. Sequence  
\_\_\_\_\_ A.A. Sequence  
\_\_\_\_\_ Structure  
\_\_\_\_\_ Bibliographic

#### Vendors

\_\_\_\_\_ IG  
\_\_\_\_\_ STN  
\_\_\_\_\_ Dialog  
\_\_\_\_\_ APS  
\_\_\_\_\_ Geninfo  
\_\_\_\_\_ SDC  
\_\_\_\_\_ DARC/Questel  
✓ \_\_\_\_\_ Other CGN

## Shears, Beverly

---

**From:** Devi, Sarvamangala  
**Sent:** Tuesday, December 09, 2003 12:46 PM  
**To:** Shears, Beverly  
**Subject:** RE: 10/878,781

Beverly:

The correct number should be 09/878,781.

-----Original Message-----

**From:** Shears, Beverly  
**Sent:** Tuesday, December 09, 2003 9:28 AM  
**To:** Devi, Sarvamangala  
**Subject:** RE: 10/878,781

Pls. re-check this number. No CRF listing avail.

-----Original Message-----

**From:** Devi, Sarvamangala  
**Sent:** Tuesday, December 09, 2003 7:37 AM  
**To:** Shears, Beverly  
**Subject:** 10/878,781

Beverly:

In application 10/878,781, would you please perform a sequence and an interference search for:

- A. A protein having the amino acid sequence of SEQ ID NO: 4;
- B. A fragment of SEQ ID NO: 4 having 1-336 amino acids; and
- C. An at least 5 amino acid-long peptide of SEQ ID NO: 4.

Thanks.

S. DEVI, Ph.D.  
AU 1645  
CM1-7E15

## Shears, Beverly

---

**From:** Devi, Sarvamangala  
**Sent:** Tuesday, December 09, 2003 7:37 AM  
**To:** Shears, Beverly  
**Subject:** 10/878,781

Beverly:

In application 10/878,781, would you please perform a sequence and an interference search for:

- A. A protein having the amino acid sequence of SEQ ID NO: 4;
- B. A fragment of SEQ ID NO: 4 having 1-336 amino acids; and
- C. An at least 5 amino acid-long peptide of SEQ ID NO: 4.

Thanks.

S. DEVI, Ph.D.  
AU 1645  
CM1-7E15

## RESULT 17

AAW55089

ID AAW55089 standard; Protein; 333 AA.

XX

AC AAW55089;

XX

DT 02-OCT-1998 (first entry)

XX

DE Streptococcus pneumoniae SP0035 protein.

XX

KW Streptococcus pneumoniae; antigen; vaccine; infection; diagnosis;  
detection; pneumonia; otitis media; meningitis.

XX

OS Streptococcus pneumoniae.

XX

PN W09818930-A2.

XX

PD 07-MAY-1998.

XX

PF 30-OCT-1997; 97WO-US19422.

XX

PR 31-OCT-1996; 96US-0029960.

XX

PA (HUMA-) HUMAN GENOME SCI INC.

XX

PI Choi GH, Hromockyj A, Johnson LS, Kunsch CA;

XX

DR WPI; 1998-272224/24.

XX

DR N-PSDB; AAV27350.

XX

PT Nucleic acid encoding antigenic peptide(s) from Streptococcus  
pneumoniae - or their epitope-containing fragments, useful in  
PT protective or therapeutic vaccines, and for diagnosis

XX

PS Claim 11; Page 59; 118pp; English.

XX

CC The present sequence represents a protein from Streptococcus pneumoniae.  
 CC The nucleic acid sequence encoding the Streptococcus pneumoniae protein  
 CC can be useful in vaccines for inducing protective antibodies against  
 CC Streptococcus pneumoniae, for treatment or prevention of infection e.g.  
 CC pneumonia, otitis media or meningitis. Probes based on the nucleic acid  
 CC are used to detect Streptococcus infection (by usual hybridisation or  
 CC amplification methods), also for isolating Streptococcus genes or their  
 CC allelic variants. The protein can be used similarly to detect specific  
 CC antibodies in standard immunoassays, especially for diagnosing or  
 CC monitoring infections. Antibodies which bind the protein are used to  
 CC detect corresponding antigens, to purify the protein and for passive  
 CC immunisation (optionally coupled to a toxin). Vaccines are administered,  
 CC e.g. by injection, orally or through the skin, typically at 0.01-1000  
 CC (especially 10-300) mu g/ml per dose.

XX

SQ Sequence 333 AA;

Query Match 17.6%; Score 59; DB 19; Length 333;

Best Local Similarity 100.0%; Pred. No. 1.3e-50;

Matches 59; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 173 GLMTTIHAYTGDQMILDGPHRGDLRRARAGAANIVPNSTGAAKAIGLVIPELNGKLDG 231

Db 171 GLMTTIHAYTGDQMILDGPHRGDLRRARAGAANIVPNSTGAAKAIGLVIPELNGKLDG 229

SEQ ID No. 4 fragment

CC antibody to provide targeting to clots. The plasmin receptor may  
CC be useful in human or veterinary medicine, for treatment of thrombosis  
CC and pulmonary embolism, and for solubilising clots in catheters or  
CC shunts.  
CC (Updated on 25-MAR-2003 to correct PF field.)  
XX  
SQ Sequence 336 AA;

Query Match 99.8%; Score 1711; DB 15; Length 336;  
Best Local Similarity 99.4%; Pred. No. 1.8e-144;  
Matches 334; Conservative 2; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 MVKVGINGRIGRLAFRIQNVGEVETRIINDLTDENMLAHLKYDTTQGRFDGTV 60  
Db 1 MVKVGINGRIGRLAFRIQNVGEVETRIINDLTDENMLAHLKYDTTQGRFDGTV 60  
Qy 61 KEGGFVNGNFIKVSARDPENIDWATDGEIVLEATGFFAKKEAAEKHLHANGAKKWI 120  
Db 61 KEGGFVNGNFIKVSARDPENIDWATDGEIVLEATGFFAKKEAAEKHLHANGAKKWI 120  
Qy 121 TAPGGNDVKTVENTNHDILDTETVISCASCTTNCLAPMAKALHDAFGIQGLMTTIIA 180  
Db 121 TAPGGNDVKTVENTNHDILDTETVISCASCTTNCLAPMAKALHDAFGIQGLMTTIIA 180  
Qy 181 YTGQMIIDGPHRGGLRRARAGAAANIVNSTGAAKAIGLVIPELNGKLDGAAQRPVPT 240  
Db 181 YTGQMIIDGPHRGGLRRARAGAAANIVNSTGAAKAIGLVIPELNGKLDGAAQRPVPT 240  
Qy 241 GSVTELVVTLDKNVSVDENAAKASNDSCFYTETDPIVSSDIVGSYGSIFDATTQKM 300  
Db 241 GSVTELVVTLDKNVSVDENAAKASNDSCFYTETDPIVSSDIVGSYGSIFDATTQKM 300  
Qy 301 EVDGSQLVKVSWYDNEMSYTAQLVRLTLEYFAKIAK 336  
Db 301 EVDGSQLVKVSWYDNEMSYTAQLVRLTLEYFAKIAK 336

See ID No. 4.

RESULT 5  
AAR56486  
ID AAR56486 standard; Protein; 336 AA.

XX AC AAR56486;

XX DT 25-MAR-2003 (updated)

DT 22-FEB-1995 (first entry)

XX DT Plasmin receptor.

XX Plasmin receptor; isolate 64/14; plasmin; alpha-2-antiplasmin;  
XX Plasminogen activator; bleeding; reocclusion; thrombosis;  
XX Pulmonary embolism; clots.

XX Streptococcus pyogenes.

XX US5328996-A.

XX 12-JUL-1994.

XX 10-AUG-1992; 92US-0928462.

XX 29-MAR-1989; 89US-0330849.

XX 16-MAY-1990; 90US-0524411.

XX 10-AUG-1992; 92US-0928462.

XX (UYFL) UNIV FLORIDA RES FOUND INC.

XX Boyle MDP, Broder C, Lottenberg R, Von Mering G;

XX WPI: 1994-225327/27.

XX N-PSDB; AAQ070705.

XX New DNA encoding bacterial plasmin receptor - useful as  
XX thrombolytic agents; used with plasminogen activator or bound to  
XX plasmin, also useful in vaccines against bacterial infection

XX Claim 1; Column 27-30; 19pp; English.

XX This sequence represents the S. pyogenes plasmin receptor. The DNA  
XX encoding this sequence was isolated from the S. pyogenes clinical  
XX isolate 64/14. The plasmin receptor has a very high affinity for  
XX plasmin which, when bound, retains its enzymatic activity but is not  
XX regulated (inhibited) by alpha-2-antiplasmin. The receptor protein,  
XX when administered concurrently or sequentially, prolongs the activity  
XX of plasminogen activator (PA) so allows a reduction in dose, and thus  
XX lowers the risk of bleeding, and may prevent reocclusion of blood  
XX vessels. The protein may be coupled to a fibrin-specific monoclonal

RESULT 1  
A42963

glyceraldehyde-3-phosphate dehydrogenase (phosphorylating) (EC 1.2.1.12) - Streptococcus  
N;Alternate names: plasmin receptor  
C;Species: Streptococcus sp.  
C;Date: 10-Mar-1994 #sequence revision 10-Mar-1994 #text\_change 03-Jun-2002  
C;Accession: A42963; B42963; JH0750  
R;Lottenberg, R.; Broder, C.C.; Boyle, M.D.; Kain, S.J.; Schroeder, B.L.; Curtiss III, J. Bacteriol. 174, 5204-5210, 1992  
A;Title: Cloning, sequence analysis, and expression in Escherichia coli of a streptococcal plasmin receptor  
A;Reference number: A42963; MUID:92355491; PMID:1322883  
A;Accession: A42963  
A;Molecule type: DNA  
A;Residues: 1-336 <LOT>  
A;Experimental source: group A, strain 64/14  
A;Note: sequence extracted from NCBI backbone (NCBIP:110308)  
A;Accession: B42963  
A;Molecule type: protein  
A;Residues: 2-74;161-164,'X',166-174;187-211,'X',213-217 <LO2>  
R;Pancholi, V.; Fischetti, V.A. J. Exp. Med. 176, 415-426, 1992  
A;Title: A major surface protein on group A streptococci is a glyceraldehyde-3-phosphate dehydrogenase  
A;Reference number: JH0750; MUID:92364544; PMID:1500854  
A;Accession: JH0750  
A;Molecule type: protein  
A;Residues: 2-30,'A',32-40 <PAN>  
C;Superfamily: glyceraldehyde-3-phosphate dehydrogenase  
C;Keywords: gluconeogenesis; glycolysis; homotetramer; NAD; oxidoreductase  
F;152/Active site: Cys #status predicted

Query Match 99.8%; Score 1711; DB 2; Length 336;  
Best Local Similarity 99.4%; Pred. No. 4e-116;  
Matches 334; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 M V V K V G I N G F G R I G R L A F R R I Q N V E G V E V T R I N D L T D P N M L A H L L K Y D T T Q G R F D G T V E V 60  
Db 1 M V V K V G I N G F G R I G R L A F R R I Q N I E G V E V T R I N D L T D P N M L A H L L K Y D T T Q G R F D G T V E V 60  
QY 61 K E G G F E V N G N F I K V S A E R D P E N I D W A T D G V E I V L E A T G F F A K K E A E K H L H A N G A K K V V I 120  
Db 61 K E G G F E V N G N F I K V S A E R D P E N I D W A T D G V E I V L E A T G F F A K K E A E K H L H A N G A K K V V I 120  
QY 121 T A P G G N D V K T V V F N T N H D I L D G T E T V I S G A S C T T N C L A P M A K A L H D A F G I Q K G L M T T I H A 180  
Db 121 T A P G G N D V K T V V F N T N H D I L D G T E T V I S G A S C T T N C L A P M A K A L H D A F G I Q K G L M T T I H A 180  
QY 181 Y T G D Q M I L D G P H R G G D L R R A G A A N I V P N S T G A A K A I G L V I P E L N G K L D G A A Q R V P V P T 240  
Db 181 Y T G D Q M I L D G P H R G G D L R R A G A A N I V P N S T G A A K A I G L V I P E L N G K L D G A A Q R V P V P T 240  
QY 241 G S V T E L V V T L D K N V S V D E I N A A M K A A S N D S F G Y T E D P I V S S D I V G V S Y G S L F D A T Q T K V M 300  
Db 241 G S V T E L V V T L D K N V S V D E I N S A M K A A S N D S F G Y T E D P I V S S D I V G V S Y G S L F D A T Q T K V M 300  
QY 301 E V D G S Q L V K V V S W Y D N E M S Y T A Q L V R T L E Y F A K I A K 336  
Db 301 E V D G S Q L V K V V S W Y D N E M S Y T A Q L V R T L E Y F A K I A K 336